

=====

Sequence Listing was accepted.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: markspencer

Timestamp: [year=2008; month=6; day=27; hr=13; min=19; sec=8; ms=610;]

=====

Application No: 10660893 Version No: 3.0

Input Set:

Output Set:

Started: 2008-06-27 11:20:24.578
Finished: 2008-06-27 11:20:25.379
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 801 ms
Total Warnings: 6
Total Errors: 0
No. of SeqIDs Defined: 6
Actual SeqID Count: 6

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)

SEQUENCE LISTING

<110> Link , Charles

<120> Methods and Compositions for Elucidating Protein Expression Profiles in Cells

<130> 05237.0003.CPU01

<140> 10660893

<141> 2003-09-12

<150> 09/811,842

<151> 2001-03-19

<150> 60/190,678

<151> 2000-03-20

<150> 60/458,152

<151> 2003-03-27

<160> 6

<170> PatentIn version 3.4

<210> 1

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> HA epitope tag

<400> 1

Tyr Pro Tyr Asp Val Pro Asp Tyr Ala

1 5

<210> 2

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> c-myc epitope tag

<400> 2

Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu

1 5 10

<210> 3

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> FLAG epitope tag

<400> 3

Asp Tyr Lys Asp Asp Asp Asp Lys

1 5

<210> 4

<211> 375

<212> DNA

<213> Artificial Sequence

<220>

<223> Gene trapped exon of HMGI-C gene

<220>

<221> misc_feature

<222> (3)..(4)

<223> n is a, c, g or t

<220>

<221> misc_feature

<222> (8)..(8)

<223> n is a, c, g or t

<220>

<221> misc_feature

<222> (350)..(350)

<223> n is a, c, g or t

<220>

<221> misc_feature

<222> (358)..(374)

<223> n is a, c, g or t

<400> 4

ttnnccgnga aagctcctcg cccttgctca ccatgggatg ccatttccta ggtctgcctc 60

ttggccgttt ttctccaatg gtctctgctt tcttctgggc tgctttagag gggctcttgt 120

ttttgctgcc tttgggtctt cctctgggtc tcttaggaga gggctcacag gttggtcttt 180

gctgctgctt cctgggtcgg ccgcgtctc gcttctgtgg caccggggcg gcaggttgtc 240

cctgggctga tgtggacggc tgcccggegc cctcaccgcg tgcgtcate ctgcctcccg 300

ccgcgctac cactgcctct cttttttttt tttttttttt tttttgaaan ccccggnnn 360

nnnnnnnnnn nnnnc 375

<210> 5

<211> 333
<212> DNA
<213> Artificial Sequence

<220>
<223> Gene trapping in pGT5A-transfected PA317 cells

<220>
<221> misc_feature
<222> (3)..(3)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (11)..(11)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (106)..(106)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (116)..(116)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (168)..(168)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (179)..(179)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (204)..(204)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (221)..(221)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (224)..(224)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (231)..(231)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (254)..(254)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (272)..(272)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (275)..(275)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (282)..(282)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (285)..(286)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (289)..(289)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (292)..(293)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (296)..(296)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (299)..(299)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (301)..(301)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (304)..(304)
<223> n is a, c, g, or t

<220>

<221> misc_feature
<222> (306)..(308)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (311)..(311)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (326)..(327)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (329)..(329)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (331)..(331)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (333)..(333)
<223> n is a, c, g, or t

<400> 5
tengcgacca nctctcgcg cttgtctacc atgggatgct cccggtggtg ggtecggtggt 60

ccctgggcag ggggtctcaa atcccgagcg agcccccaaa tgaaanaccc ccgtcntggg 120

tagtcaatca ctgagaggag accctcccaa ggaacagcga gacctntt cggtatgcana 180

cagcaagagg ctttattggg aatnccggta cccgggcgac ncantctatc ngaagactgg 240

cgttatTTTT tttntTTTT ttttttgaat tncngggac anccnnctna gnntanctnc 300

nctntnnnct nccctcctta cttctnntnt ntn 333

<210> 6
<211> 11
<212> DNA
<213> Artificial Sequence

<220>
<223> pGT-fs2

<400> 6
gagtcaccagc t